#5

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PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants:	Feighner, Scott D. *et al.*		
Serial No.:	09/719,485	Case No.: 20251P	Art Unit: Unassigned
Filed:	December 12, 2000		Examiner: Unassigned
For:	CLONING AND IDENTIFICATION OF THE MOTILIN RECEPTOR		

Assistant Commissioner for Patents
Washington, D.C. 20231

RESPONSE TO COMPLY WITH SEQUENCE REQUIREMENT

Sir:

The following statements are made as required by 37 C.F.R. § 1.821(f).

I hereby state that the contents of the paper and computer readable copies of the Sequence Listing, submitted in accordance with 37 C.F.R. § 1.821 (c) and (e), respectively, are the same and contain no new matter.

I hereby certify that this correspondence is being deposited with the United States Postal Service as first class mail in an envelope addressed to: Assistant Commissioner for Patents, Washington, D.C. 20231, on the date appearing below.

MERCK & CO., INC.

By _____ Date 5/23/2001

Respectfully submitted,

By _____

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Reg. No. 42,452
Attorney for Applicant(s)

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Date: May 23, 2001

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MacNeil, Douglas
Howard, Andrew D.
Pong, Sheng-Shung
Smith, Roy G.

<120> CLONING AND IDENTIFICATION OF THE MOTILIN RECEPTOR

<130> 20251P

<140> 09/719,485
<141> 2000-12-12

<150> PCT/US99/12773
<151> 1999-06-08

<150> 60/089,098
<151> 1998-06-12

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<212> PRT
<213> Homo sapiens

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        35                  40                  45
Phe Val Val Gly Val Ser Gly Asn Val Val Thr Val Met Leu Ile Gly
    50                  55                  60
Arg Tyr Arg Asp Met Arg Thr Thr Thr Asn Leu Tyr Leu Gly Ser Met
65                  70                  75                  80
Ala Val Ser Asp Leu Leu Ile Leu Leu Gly Leu Pro Phe Asp Leu Tyr
                85                  90                  95
Arg Leu Trp Arg Ser Arg Pro Trp Val Phe Gly Pro Leu Leu Cys Arg
            100                 105                 110
Leu Ser Leu Tyr Val Gly Glu Gly Cys Thr Tyr Ala Thr Leu Leu His
        115                 120                 125
Met Thr Ala Leu Ser Val Glu Arg Tyr Leu Ala Ile Cys Arg Pro Leu
    130                 135                 140
Arg Ala Arg Val Leu Val Thr Arg Arg Arg Val Arg Ala Leu Ile Ala
145                 150                 155                 160
Val Leu Trp Ala Val Ala Leu Leu Ser Ala Gly Pro Phe Leu Phe Leu
                165                 170                 175
Val Gly Val Glu Gln Asp Pro Gly Ile Ser Val Val Pro Gly Leu Asn
            180                 185                 190
Gly Thr Ala Arg Ile Ala Ser Ser Pro Leu Ala Ser Ser Pro Pro Leu
        195                 200                 205
Trp Leu Ser Arg Ala Pro Pro Pro Ser Pro Pro Ser Gly Pro Glu Thr
    210                 215                 220
Ala Glu Ala Ala Ala Leu Phe Ser Arg Glu Cys Arg Pro Ser Pro Ala
225                 230                 235                 240
Gln Leu Gly Ala Leu Arg Val Met Leu Trp Val Thr Thr Ala Tyr Phe
                245                 250                 255
Phe Leu Pro Phe Leu Cys Leu Ser Ile Leu Tyr Gly Leu Ile Gly Arg
            260                 265                 270
Glu Leu Trp Ser Ser Arg Arg Pro Leu Arg Gly Pro Ala Ala Ser Gly
        275                 280                 285
Arg Glu Arg Gly His Arg Gln Thr Val Arg Val Leu Leu Val Val Val
    290                 295                 300
Leu Ala Phe Ile Ile Cys Trp Leu Pro Phe His Val Gly Arg Ile Ile
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Tyr Ile Asn Thr Glu Asp Ser Arg Met Met Tyr Phe Ser Gln Tyr Phe
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Leu Leu Leu Ala Arg Lys Ser Arg Pro Arg Gly Phe His Arg Ser Arg
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gccgtgtccg acctactcat cctgctcggg ctgccgttcg acctgtaccg cctctggcgc     300
tcgcggccct gggtgttcgg gccgctgctc tgccgcctgt ccctctacgt gggcgagggc     360
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<212> PRT
<213> Homo sapiens

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Phe Val Val Gly Val Ser Gly Asn Val Val Thr Val Met Leu Ile Gly
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                85                  90                  95
Arg Leu Trp Arg Ser Arg Pro Trp Val Phe Gly Pro Leu Leu Cys Arg
            100                 105                 110
Leu Ser Leu Tyr Val Gly Glu Gly Cys Thr Tyr Ala Thr Leu Leu His
        115                 120                 125
Met Thr Ala Leu Ser Val Glu Arg Tyr Leu Ala Ile Cys Arg Pro Leu
    130                 135                 140
Arg Ala Arg Val Leu Val Thr Arg Arg Arg Val Arg Ala Leu Ile Ala
145                 150                 155                 160
Val Leu Trp Ala Val Ala Leu Leu Ser Ala Gly Pro Phe Leu Phe Leu
                165                 170                 175
Val Gly Val Glu Gln Asp Pro Gly Ile Ser Val Val Pro Gly Leu Asn
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Gly Thr Ala Arg Ile Ala Ser Ser Pro Leu Ala Ser Ser Pro Pro Leu
        195                 200                 205
Trp Leu Ser Arg Ala Pro Pro Pro Ser Pro Pro Ser Gly Pro Glu Thr
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Ala Glu Ala Ala Ala Leu Phe Ser Arg Glu Cys Arg Pro Ser Pro Ala
225                 230                 235                 240
Gln Leu Gly Ala Leu Arg Val Met Leu Trp Val Thr Thr Ala Tyr Phe
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Glu Leu Trp Ser Ser Arg Arg Pro Leu Arg Gly Pro Ala Ala Ser Gly
        275                 280                 285
Arg Glu Arg Gly His Arg Gln Thr Val Arg Val Leu Arg Lys Trp Ser
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Gly Gly Ser Gly Ile Tyr Asn Leu Leu Val Ala Leu Pro Arg Trp Gln
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<212> PRT
<213> Spheroides nephelus

<400> 7

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Phe Pro Ala Ser Thr Leu Ile Pro Val Thr Val Ile Cys Ile Leu Ile
        35                  40                  45
Phe Val Val Gly Val Thr Gly Asn Thr Met Thr Ile Leu Ile Ile Gln
    50                  55                  60
Tyr Phe Lys Asp Met Lys Thr Thr Thr Asn Leu Tyr Leu Ser Ser Met
65                  70                  75                  80
Ala Val Ser Asp Leu Val Ile Phe Leu Cys Leu Pro Phe Asp Leu Tyr
                85                  90                  95
Arg Leu Trp Lys Tyr Val Pro Trp Leu Phe Gly Glu Ala Val Cys Arg
            100                 105                 110
Leu Tyr His Tyr Ile Phe Glu Gly Cys Thr Ser Ala Thr Ile Leu His
        115                 120                 125
Ile Thr Ala Leu Ser Ile Glu Arg Tyr Leu Ala Ile Ser Phe Pro Leu
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Arg Ser Lys Val Met Val Thr Arg Arg Arg Val Gln Tyr Ile Ile Leu
145                 150                 155                 160
Ala Leu Trp Cys Phe Ala Leu Val Ser Ala Ala Pro Thr Leu Phe Leu
                165                 170                 175
Val Gly Val Glu Tyr Asp Asn Glu Thr His Pro Asp Tyr Asn Thr Gly
            180                 185                 190
Gln Cys Lys His Thr Gly Tyr Ala Ile Ser Ser Gly Gln Leu His Ile
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Met Ile Trp Val Ser Thr Thr Tyr Phe Phe Cys Pro Met Leu Cys Leu
    210                 215                 220
Leu Phe Leu Tyr Gly Ser Ile Gly Cys Lys Leu Trp Lys Ser Lys Asn
225                 230                 235                 240
Asp Leu Gln Gly Pro Cys Ala Leu Ala Arg Glu Arg Ser His Arg Gln
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Thr Val Lys Ile Leu Val Val Val Val Leu Ala Phe Ile Ile Cys Trp
            260                 265                 270
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Asp Thr Ala Met Leu Ser Gln Asn Phe Asn Met Ala Ser Met Val Leu
290 295 300
Cys Tyr Leu Ser Ala Ser Ile Asn Pro Val Val Tyr Asn Leu Met Ser
305 310 315 320
Arg Lys Tyr Arg Ala Ala Ala Lys Arg Leu Phe Leu Leu His Gln Arg
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<213> Artificial Sequence

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<223> PCR Primer

<400> 8
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<212> DNA
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<212> DNA
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<211> 300
<212> PRT
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Phe Pro Leu Gly Ala Leu Val Pro Val Thr Ala Val Cys Leu Cys Leu
35 40 45
Phe Val Val Gly Val Ser Gly Asn Val Val Thr Val Met Leu Ile Gly
50 55 60
Arg Tyr Arg Asp Met Arg Thr Thr Thr Asn Leu Tyr Leu Gly Ser Met
65 70 75 80
Ala Val Ser Asp Leu Leu Ile Leu Leu Gly Leu Pro Phe Asp Leu Tyr
85 90 95
Arg Leu Trp Arg Ser Arg Pro Trp Val Phe Gly Pro Leu Leu Cys Arg
100 105 110
Leu Ser Leu Tyr Val Gly Glu Gly Cys Thr Tyr Ala Thr Leu Leu His
115 120 125
Met Thr Ala Leu Ser Val Glu Arg Tyr Leu Ala Ile Cys Arg Pro Leu
130 135 140
Arg Ala Arg Val Leu Val Thr Arg Arg Arg Val Arg Ala Leu Ile Ala
145 150 155 160
Val Leu Trp Ala Val Ala Leu Leu Ser Ala Gly Pro Phe Leu Phe Leu
165 170 175
Val Gly Val Glu Gln Asp Pro Gly Ile Ser Val Val Pro Gly Leu Asn
180 185 190
Gly Thr Ala Arg Ile Ala Ser Ser Pro Leu Ala Ser Ser Pro Pro Leu
195 200 205
Trp Leu Ser Arg Ala Pro Pro Pro Ser Pro Pro Ser Gly Pro Glu Thr
210 215 220
Ala Glu Ala Ala Ala Leu Phe Ser Arg Glu Cys Arg Pro Ser Pro Ala
225 230 235 240
Gln Leu Gly Ala Leu Arg Val Met Leu Trp Val Thr Thr Ala Tyr Phe
245 250 255
Phe Leu Pro Phe Leu Cys Leu Ser Ile Leu Tyr Gly Leu Ile Gly Arg
260 265 270
Glu Leu Trp Ser Ser Arg Arg Pro Leu Arg Gly Pro Ala Ala Ser Gly
275 280 285

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<213> Homo sapiens

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tttcctattt cgattccagc ctccacccgc cggt 154

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<211> 602
<212> DNA
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ctctgcaact tttctatctg agcgcatcta tcaacccaat cctctacaac ctcatttcaa 180
agaagtacag agcggcggcc tttaaactgc tgctcgcaag gaagtccagg ccgagaggct 240
tccacagaag cagggacact gcgggggaag ttgcagggga cactggagga gacacggtgg 300
gctacaccga gacaagcgct aacgtgaaga cgatgggata acgtaagtgg agccgccgtg 360
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aa 602

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<212> PRT
<213> Homo sapiens

<400> 15

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Gly Arg Ile Ile Tyr Ile Asn Thr Glu Asp Ser Arg Met Met Tyr Phe
            20                  25                  30
Ser Gln Tyr Phe Asn Ile Val Ala Leu Gln Leu Phe Tyr Leu Ser Ala
        35                  40                  45
Ser Ile Asn Pro Ile Leu Tyr Asn Leu Ile Ser Lys Lys Tyr Arg Ala
    50                  55                  60
Ala Ala Phe Lys Leu Leu Leu Ala Arg Lys Ser Arg Pro Arg Gly Phe
65                  70                  75                  80
His Arg Ser Arg Asp Thr Ala Gly Glu Val Ala Gly Asp Thr Gly Gly
                85                  90                  95
Asp Thr Val Gly Tyr Thr Glu Thr Ser Ala Asn Val Lys Thr Met Gly
            100                 105                 110
Arg Lys Trp Ser Arg Arg Gly Ser Lys Asp Ala Cys Leu Gln Ser Ala
        115                 120                 125
Pro Pro Gly Thr Ala Gln Thr Leu Gly Pro Leu Pro Leu Leu Ala Gln
    130                 135                 140
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Leu Trp Ala Pro Leu Pro Ala Pro Phe Pro Ile Ser Ile Pro Ala Ser
145                 150                 155                 160
Thr Arg Arg Gly Gly Gly Ser Gly Ile Tyr Asn Leu Leu Val Ala Leu
                165                 170                 175
Pro Arg Trp Gln Asn His Leu His Lys His Gly Arg Phe Ala Asp Asp
            180                 185                 190
Val Leu Leu Ser Val Leu
        195
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